



SEQUENCE LISTING

<110> Harberd, Nicholas P
Richards, Donald E
Peng, Jinrong

<120> Genetic Control of Plant Growth and Development

<130> 620-298

<140> US 10/809,945

<141> 2004-03-26

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<151> 2000-03-01

<150> PCT/GB98/02383

<151> 1998-08-07

<150> GB 9717192.0

<151> 1997-08-13

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<170> PatentIn Ver. 2.0

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Ala Gln Lys Leu Glu Lys Leu Glu Met Ala Met Gly Met Gly Gly Val
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Asp Thr Val Xaa Tyr Asn Pro Thr Asp Xaa Ser Ser Trp Val Glu Ser
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165 170 175

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195 200 205

Val Glu Ala Ala Pro Pro Val Ala Ala Ala Ala Asn Ala Thr Pro Ala
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Leu Pro Val Val Val Val Asp Thr Gln Glu Ala Gly Ile Arg Leu Val
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 65 70 75 80
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 Ala Ala Ala Asp Ser Ser Ser Ser Thr Tyr Ala Leu Arg Pro Ile Ser
 145 150 155 160
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 Asp Thr Lys Arg Met Arg Thr Gly Gly Gly Ser Thr Ser Ser Ser Ser
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Tyr Glu Ser Cys Pro Tyr Leu Lys Phe Ala His Phe Thr Ala Asn Gln
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Lys Val Leu Gly Thr Val Arg Ala Val Arg Pro Arg Ile Val Thr Val
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Glu Arg His Glu Thr Leu Gly Gln Trp Arg Asn Arg Leu Gly Asn Ala
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Glu Asn Phe Ser Ala Ala Glu Ala Leu Val Lys Gln Ile Pro Met Leu
260 265 270
Ala Ser Ser Gln Gly Gly Ala Met Arg Lys Val Ala Ala Tyr Phe Gly

275	280	285
Glu Ala Leu Ala Arg Arg Val Tyr Arg Phe Arg Pro Pro Pro Asp Ser		
290	295	300
Ser Leu Leu Asp Ala Ala Phe Ala Asp Leu Leu His Ala His Phe Tyr		
305	310	315 320
Glu Ser Cys Pro Tyr Leu Lys Phe Ala His Phe Thr Ala Asn Gln Ala		
	325	330 335
Ile Leu Glu Ala Phe Ala Gly Cys Arg Arg Val His Val Val Asp Phe		
	340	345 350
Gly Ile Lys Gln Gly Met Gln Trp Pro Ala Leu Leu Gln Ala Leu Ala		
	355	360 365
Leu Arg Pro Gly Gly Pro Pro Ser Phe Arg Leu Thr Gly Val Gly Pro		
	370	375 380
Pro Gln Pro Asp Glu Thr Asp Ala Leu Gln Gln Val Gly Trp Lys Leu		
	385	390 395 400
Ala Gln Phe Ala His Thr Ile Arg Val Asp Phe Gln Tyr Arg Gly Leu		
	405	410 415
Val Ala Ala Thr Leu Ala Asp Leu Glu Pro Phe Met Leu Gln Pro Glu		
	420	425 430
Gly Asp Asp Thr Asp Asp Glu Pro Glu Val Ile Ala Val Asn Ser Val		
	435	440 445
Phe Glu Leu His Arg Leu Leu Ala Gln Pro Gly Ala Leu Glu Lys Val		
	450	455 460
Leu Gly Thr Val Arg Ala Val Arg Pro Arg Ile Val Thr Val Val Glu		
	465	470 475 480
Gln Glu Ala Asn His Asn Ser Gly Thr Phe Leu Asp Arg Phe Thr Glu		
	485	490 495
Ser Leu His Tyr Tyr Ser Thr Met Phe Asp Ser Leu Glu Gly Ala Gly		
	500	505 510
Ala Gly Ser Gly Gln Ser Thr Asp Ala Ser Pro Ala Ala Ala Gly Gly		
	515	520 525
Thr Asp Gln Val Met Ser Glu Val Tyr Leu Gly Arg Gln Ile Cys Asn		
	530	535 540
Val Val Ala Cys Glu Gly Ala Glu Arg Thr Glu Arg His Glu Thr Leu		
	545	550 555 560
Gly Gln Trp Arg Ser Arg Leu Gly Gly Ser Gly Phe Ala Pro Val His		
	565	570 575
Leu Gly Ser Asn Ala Tyr Lys Gln Ala Ser Thr Leu Leu Ala Leu Phe		

580

585

590

Ala Gly Gly Asp Gly Tyr Arg Val Glu Glu Lys Asp Gly Cys Leu Thr
 595 600 605

Leu Gly Trp His Thr Arg Pro Leu Ile Ala Thr Ser Ala Trp Arg Val
 610 615 620

Ala Ala Ala Ala Ala Pro
 625 630

<210> 9

<211> 100

<212> PRT

<213> Zea mays

<400> 9

Tyr Gln Asp Ala Gly Gly Ser Gly Gly Asp Met Gly Ser Ser Lys Asp
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Lys Met Met Ala Ala Ala Ala Gly Ala Gly Glu Gln Glu Glu Glu Asp
 20 25 30

Val Asp Glu Leu Leu Ala Ala Leu Gly Tyr Lys Val Arg Ser Ser Asp
 35 40 45

Met Ala Gly Leu Glu Gln Leu Glu Met Ala Met Gly Met Gly Gly Val
 50 55 60

Gly Gly Ala Gly Ala Thr Ala Asp Asp Gly Phe Val Ser His Leu Ala
 65 70 75 80

Thr Asp Thr Val His Tyr Asn Pro Ser Asp Leu Ser Ser Trp Val Glu
 85 90 95

Ser Met Leu Ser
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<210> 10

<211> 123

<212> PRT

<213> Zea mays

<400> 10

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 1 5 10 15

Glu Glu Glu Asp Val Asp Glu Leu Leu Ala Ala Leu Gly Tyr Lys Val
 20 25 30

Arg Ser Ser Asp Met Ala Asp Val Ala Gln Lys Leu Glu Gln Leu Glu
 35 40 45

Met Ala Met Gly Met Gly Gly Val Gly Gly Ala Gly Ala Thr Ala Asp

- 50

55

60

Asp Gly Phe Val Ser His Leu Ser Ser Trp Val Glu Ser Met Leu Ser
65 70 75 80
Glu Leu Asn Ala Pro Pro Ala Pro Leu Pro Pro Ala Thr Pro Ala Pro
85 90 95
Arg Leu Ala Ser Thr Ser Ser Thr Val Thr Ser Gly Ala Ala Ala Gly
100 105 110
Ala Gly Tyr Phe Asp Leu Pro Pro Ala Val Asp
115 120

<210> 11
<211> 138
<212> PRT
<213> Triticum aestivum

<400> 11
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Gln Lys Leu Glu Gln Leu Glu Met Ala Met Gly Met Gly Gly Val Gly
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Ala Gly Ala Ala Pro Asp Asp Ser Phe Ala Thr His Leu Ala Thr Asp
35 40 45
Thr Val His Tyr Asn Pro Thr Asp Leu Ser Ser Trp Val Glu Ser Met
50 55 60
Leu Ser Glu Leu Asn Ala Ser Thr Ser Ser Thr Val Thr Gly Ser Gly
65 70 75 80
Gly Tyr Phe Asp Leu Pro Pro Ser Val Asp Ser Ser Ser Ser Ile Tyr
85 90 95
Ala Leu Arg Pro Ile Pro Ser Pro Ala Gly Ala Thr Ala Pro Ala Asp
100 105 110
Leu Ser Ala Asp Ser Val Arg Asp Pro Lys Arg Met Arg Thr Gly Gly
115 120 125
Ser Ser Thr Ser Ser Ser Ser Ser Ser Ser
130 135

<210> 12
<211> 770
<212> DNA
<213> Oryza sativa

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tgctggcggc	gctcgggtac	aaggtgcggt	cgcccgacat	ggccgacgtc	gcgcagaagc	180
tggagcagct	ggagatggcc	atggggatgg	gcggcgtgag	cgcccccgcc	gccgcggatg	240
acgggttcgt	gtcgcacctg	gccacggaca	ccgtgcacta	caaccctctg	gacctctcct	300
cctgggtcga	gagcatgctt	tccgagctca	acgcgcgcgt	gccccctatc	ccgccagcgc	360
cgccggctgc	ccgccatgct	tccacctcgt	ccactgtcac	cggcggcggt	ggtagcggct	420
tctttgaact	cccagccgct	gccgactcgt	cgagtagcac	ctacgccctc	aggccgatct	480
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cggcgaatgc	gcccgcgcgt	ccggttggtg	tggttgacac	gcaggaggct	gggatccggc	720
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<210> 13

<211> 1768

<212> DNA

<213> Triticum aestivum

<400> 13

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ccgcctccat	gatcaccggt	aaaactccta	agccctatta	ttactactat	tatgtttaaa	1620
tgtctattat	tgctatgtgt	aattcctcca	accgctcata	tcaaaaataag	cacgggcccgg	1680
aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	aaaaaaaaaa	1740
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<210> 14

<211> 2125

<212> DNA

<213> Triticum aestivum

<400> 14

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cagccggaca	gctccctcct	cgacgcgcgc	ttcgccgacc	tcctccacgc	gcacttctac	960
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<210> 15

<211> 2255

<212> DNA

<213> Zea mays

<400> 15

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 <211> 302
 <212> DNA
 <213> Zea mays

<400> 16						
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ga						302

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 <211> 371
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 <213> Zea mays

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gagctcaacg	cgccccagc	gcccgtcccg	cccgcgacgc	cgcccccaag	gctcgcgtcc	300
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<210> 18
 <211> 416
 <212> DNA
 <213> Triticum aestivum

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atccccctccc cggccggcgc gacggcgccg gccgacctgt ccgccgactc cgtgcgggat 360
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<210> 19

<211> 725

<212> DNA

<213> Oryza sativa

<220>

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<222> (171)

<223> n is any nucleotide

<220>

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<222> (302)

<223> n is any nucleotide

<220>

<221> misc_feature

<222> (427)

<223> n is any nucleotide

<220>

<221> misc_feature

<222> (444)

<223> n is any nucleotide

<220>

<221> misc_feature

<222> (459)

<223> n is any nucleotide

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<221> misc_feature

<222> (711)

<223> n is any nucleotide

<400> 19

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 ggtgc 725

<210> 20
 <211> 258
 <212> PRT
 <213> Oryza sativa

<220>
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 <223> Xaa is unknown or other amino acid

<220>
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 <222> (143)
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<220>
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 <222> (148)
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<220>
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 <222> (250)
 <223> Xaa is unknown or other amino acid

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 Glu Asp Val Asp Glu Leu Leu Ala Ala Leu Gly Tyr Lys Val Arg Ser
 35 40 45
 Ser Asp Met Ala Asp Val Ala Gln Xaa Leu Glu Gln Leu Glu Met Ala
 50 55 60
 Met Gly Met Gly Gly Val Ser Ala Pro Gly Ala Ala Asp Asp Gly Phe
 65 70 75 80
 Val Ser His Leu Ala Thr Asp Thr Val His Tyr Asn Pro Ser Asp Leu
 85 90 95
 Ser Ser Trp Val Glu Ser Met Leu Ser Glu Leu Lys Ala Pro Leu Pro
 100 105 110
 Leu Ile Pro Pro Gly Ala Ala Gly Leu Pro Ala Met Leu Ser Pro Thr
 115 120 125
 Ser Ser Thr Val Thr Gly Gly Gly Gly Ser Gly Phe Phe Glu Xaa Pro

130	135	140
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145	150	155 160
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	165	170 175
Asp Thr Lys Arg Met Arg Thr Gly Gly Gly Ser Thr Ser Ser Ser Ser		
	180	185 190
Ser Ser Ser Ser Ser Leu Gly Gly Gly Ala Ser Arg Gly Ser Val Val		
	195	200 205
Glu Ala Ala Pro Pro Ala Thr Gln Gly Ala Ala Ala Ala Asn Ala Pro		
	210	215 220
Ala Val Pro Val Val Val Val Asp Thr Gln Glu Glu Glu Ala Gly Ile		
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Arg Leu Val His Ala Leu Leu Ala Cys Xaa Glu Ala Val Gln Gln Glu		
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Asn Phe

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gcactttctac nagtcttgcc cctacctcaa gttcgcgcac ttcaccgcca attaggccat 180
cctggaggcg ttgcccggct gccgccgct gcacgtcgtc gacttcggca tcaagcaggg 240
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ggaggccggg attcggntgg tncacgcgct gctggngtgc gnggagnccg tgcagcagga 180
gaacctctcc gccgcggagg cgctngtgaa gnagataccc ntgctggccc agtcccaggg 240
cggcgagatg ngcaaggtn gacgttactt ngnagangcc ctgcccgcg gagtgattcc 300
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tacctcggcc ggcagatctg caacgtggtg gctgcgagg gggcggagcg cacantancg 240
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gtcttgggtc gagagcatgc tgtcggagct aaangagccg cngccgcccc tccgcccgc 240
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ttggccagag atagatagag aggcgaggtta gctcgcggat catgaagcgg gaggaccagg 180
acgccggagg gagcggcggc ggcggtggcg gcatgggttc gtccgaggac aagatgatgg 240
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aggtgcgcgc ctccgacatg gcggacgtgg cgcagaagct ggagcagctc gagatggcca 360
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acgtccaaaa ctgggctgag ctttgttctg ttctgttatg ttcattggtgc tcaactgctct 300
gatgaacatg atggtgcctc caatggtggc tttgcaattg ttgaaacgtt tggcttgggg 360
gacttgngtg ggtgggtgca tggggatgaa tattcacatc nccggattaa aattaagcca 420
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tcggcggcgg  cggggggacgg  ggaggaggtg  cacaacnttt  nggcgggact  cgngtaccac  180
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tcgccaccca cctcgccacg gacaccggca cacaaccca ccgacctgtc gtcttggggtc 180
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cgccgccttc gccgacctcc tccacgcgca cttctacgag tctgcccct acctcaagtt 180
cgcgcacttc accgccaacc aggccatcct ggaggcggtc gccggctgcc gccgcgtgca 240
cgtcgtcgac ttcggcatca agcaggggat gcagtggccc gcacttctcc aggccctcgc 300
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cgggccggcg cgacggcgcc ggccgacctg tccgccgact ccgtgcggga tcccaagcgg 180
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tctccggcga acgcacgcac gcacgcactt gaagaagaag aagctaaatg tcatgtcagt 180
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taatcccttc ttcgttccca gtttctccac cgctccatg atcaccccggt aaaactccta 360
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nacaggtcgg tggggttgta gtgcacggtg tccgtggcga ggggggtggcn aanctgtcgt 180
caggggcggc gccngcgccc acnccgccc tcccatggc catctcganc tgctccagct 240
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<220>
<221> misc_feature
<222> (385)
<223> n is any nucleotide

<400> 77
gaccaccacc ggcattgtagt aatgtaatcc cttcttcntt cccagttctc caccgcctcc 60
atgatcacc gtaaaactcc taagccctat tattactact attatgtnta aatgtctatt 120
attgctangt gtaattcctc caaccgctca tatcaaaata agcacgggcc ggactttgtt 180
agcagctcca atgagaatga aatgaatttt gtaacgcaagg cacgtccaaa actgggctga 240
gctttgttct gttctgttat gttcatgggtg ctcaactgctc tgatgaacat gatgggtgct 300
ccaatgggtg gctttgcaat tgttgaacgt tttggcttgg gggacttggt gnntgggtgca 360
tgggaatgaa nattccacat ccncnggaat taaaattagc ccatcccg 408

<210> 78
<211> 84
<212> PRT
<213> Arabidopsis thaliana

<400> 78
Met Lys Arg Asp His His His His Gln Asp Lys Lys Thr Met Met
1 5 10 15
Met Asn Glu Glu Asp Asp Gly Asn Gly Met Asp Glu Leu Leu Ala Val
20 25 30
Leu Gly Tyr Lys Val Arg Ser Ser Glu Met Ala Asp Val Ala Gln Lys
35 40 45
Leu Glu Gln Leu Glu Val Met Met Ser Asn Val Gln Glu Asp Asp Leu
50 55 60
Ser Gln Leu Ala Thr Glu Thr Val His Tyr Asn Pro Ala Glu Leu Tyr
65 70 75 80
Thr Trp Leu Asp

<210> 79
<211> 87
<212> PRT
<213> Oryza sativa

<220>
<221> SITE
<222> (26)
<223> Xaa is unknown or other amino acid

<220>
<221> SITE
<222> (31)
<223> Xaa is unknown or other amino acid

<400> 79
Glu Ala Gly Gly Ser Ser Gly Gly Gly Ser Ser Ala Asp Met Gly Ser
1 5 10 15
Cys Lys Asp Lys Val Met Ala Gly Ala Xaa Gly Glu Glu Glu Xaa Val
20 25 30
Asp Glu Leu Leu Ala Ala Leu Gly Tyr Lys Val Arg Ser Ser Asp Met
35 40 45
Ala Asp Val Ala Gln Lys Leu Glu Gln Leu Glu Met Ala Met Gly Met
50 55 60
Gly Gly Val Thr Pro Pro Ala Gln Arg Met Thr Gly Ser Cys Arg Thr
65 70 75 80
Trp Pro Arg Thr Lys Phe Ile
85

<210> 80
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 80
ggc gatgaca cggatgacg 19

<210> 81
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 81
cttg cgc atg gcacc gccct gcgac gaag 29

<210> 82
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 82
 ccagctaata atggcttgcg cgcctcg 27

<210> 83
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 83
 tatcccagaa ccgaaaccga g 21

<210> 84
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 84
 cggcgtcttg gtactcgcg ttcattg 26

<210> 85
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 85
 tgggctcccg cgccgagtcc gtggac 26

<210> 86
 <211> 31
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 86
 ctccaagcct cttgcgctga ccgagatcga g 31

<210> 87
 <211> 31
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Primer

<400> 87
 tccacaggct caccagtcac caacatcaat c 31

<210> 88
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 88
acggtactgg aagtccacgc ggatggtgtg 30

<210> 89
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 89
cgcacaccat ccgctggac ttccagtac 29

<210> 90
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 90
ctcggccggc agatctgcaa cgtggtg 27

<210> 91
<211> 33
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 91
ttgtgacggt ggacgatgtg gacgcgagcc ttg 33

<210> 92
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 92
ggacgtgcg acaaaccgtc catcgatcca ac 32

<210> 93

<211> 30
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Primer

 <400> 93
 tccgaaatca tgaagcgcgga gtaccaagac 30

 <210> 94
 <211> 29
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Primer

 <400> 94
 tcgggtacaa ggtgcgttcg tcggatatg 29

 <210> 95
 <211> 21
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Primer

 <400> 95
 atgaagcgcg agtaccaaga c 21

 <210> 96
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Primer

 <400> 96
 gtgtgccttg atgcggtcca gaag 24

 <210> 97
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence: Primer

 <400> 97
 aaccaccct ccctgatcac ggag 24

 <210> 98
 <211> 26
 <212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 98

cactaggagc tccgtggctg aagctg

26

<210> 99

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 99

gctgcgcaag aagccggtgc agctc

25

<210> 100

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 100

agtacacttc cgacatgact tg

22

<210> 101

<211> 4

<212> PRT

<213> Zea mays

<400> 101

Val Ala Gln Lys

1

<210> 102

<211> 12

<212> PRT

<213> Zea mays

<400> 102

Leu Ala Thr Asp Thr Val His Tyr Asn Pro Ser Asp

1

5

10

<210> 103

<211> 13

<212> PRT

<213> Triticum aestivum

<400> 103

Leu Asn Ala Pro Pro Pro Pro Leu Pro Pro Ala Pro Gln
1 5 10

<210> 104
<211> 17
<212> PRT
<213> Triticum aestivum

<400> 104
Asp Glu Leu Leu Ala Ala Leu Gly Tyr Lys Val Arg Ala Ser Asp Met
1 5 10 15

Ala

<210> 105
<211> 51
<212> DNA
<213> Triticum aestivum

<400> 105
gacgagctgc tggcggcgct cgggtacaag gtgcgcgcct ccgacatggc g 51

<210> 106
<211> 17
<212> PRT
<213> Zea mays

<400> 106
Asp Glu Leu Leu Ala Ala Leu Gly Tyr Lys Val Arg Ser Ser Asp Met
1 5 10 15

Ala

<210> 107
<211> 5
<212> PRT
<213> Arabidopsis thaliana

<400> 107
Asp Glu Leu Leu Ala
1 5

<210> 108
<211> 4
<212> PRT
<213> Arabidopsis thaliana

<400> 108
Glu Gln Leu Glu
1